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Figure 1-A

>1737775	GAAATCACAG	GGAGATGTAC	AGCAATGGGG	CCATTTAAGA	GTTCTGTGTT
>1737775IH	GAAATCACAG	GGAGATGTAC	AGCAATGGGG	CCATTTAAGA	GTTCTGTGTT
Consensus	GAAATCACAG	GGAGATGTAC	AGCAATGGGG	CCATTTAAGA	GTTCTGTGTT
			·		
>1737775	CATCTTGATT	CTTCACCTTC	TAGAAGGGGC	CCTGAGTAAT	TCACTCATTC
>1737775IH	CATCTTGATT	CTTCACCTTC	TAGAAGGGGC	CCTGAGTAAT	TCACTCATTC
Consensus	CATCTTGATT	CTTCACCTTC	TAGAAGGGGC	CCTGAGTAAT	TCACTCATTC
>1737775	AGCTGAACAA	CAATGGCTAT	GAAGGCATTG	TCGTTGCAAT	CGACCCCAAT
>1737775IH	AGCTGAACAA	CAATGGCTAT	GAAGGCATTG	TCGTTGCAAT	CGACCCCAAT
>776455				GTTGCAAT	CGACCCCAAT
Consensus	AGCTGAACAA	CAATGGCTAT	GAAGGCATTG	TCGTTGCAAT	CGACCCCAAT
>1737775	GTGCCAGAAG	ATGAAACACT	CATTCAACAA	ATAAAGGACA	TGGTGACCCA
>1737775IH	GTGCCAGAAG	ATGAAACACT	CATTCAACAA	ATAAAGGACA	TGGTGACCCA
>776455	GTGCCAGAAG	ATGAAACACT	CATTCAACAA	ATAAAGGACA	TGGTGACCCA
Consensus	GTGCCAGAAG	ATGAAACACT	CATTCAACAA	ATAAAGGACA	TGGTGACCCA
>1737775	GGCATCTCTG	TATCTGTTTG	AAG		
>1737775IH	GGCATCTCTG	TATCTGTTTG	AAGCTACAGG	AAAGCGATTT	TATTTCAAAA
>776455	GGCATCTCTG	TATCTGTTTG	AAGCTACAGG	AAAGCGATTT	TATTTCAAAA
Consensus	GGCATCTCTG	TATCTGTTTG	AAGCTACAGG	AAAGCGATTT	TATTTCAAAA
>1737775IH	ATGTTGCCAT	TTTGATTCCT	GAAACATGGA	AGACAAAGGC	TGACTATGTG
>776455	ATGTTGCCAT	TTTGATTCCT	GAAACATGGA	AGACAAAGGN	TGACTATGTG
Consensus	ATGTTGCCAT	TTTGATTCCT	GAAACATGGA	AGACAAAGGC	TGACTATGTG
>1737775IH	AGACCAAAAC	TTGAGACCTA	CAAAAATGCT	GATGTTCTGG	TTGCTGAGTC
>776455	AGACCAAAAC	TTGAGACCTA	CAAAAATGCT	GATGTTCTGG	TTGCTGAGTC
>3234527			AAAATGCT	GATGTTCTGG	TTGCTGAGTC
Consensus	AGACCAAAAC	TTGAGACCTA	CAAAAATGCT	GATGTTCTGG	TTGCTGAGTC
		•			
>1737775IH	TACTCCTCCA	GGTAATGATG	AACCCTACAC	TGAGCAGATG	GGCAACTGTG
>776455	TANTCCTCCA	GGNAATGATG	AACCCTACAC	TGNGCAGATG	GGCAACTGTG
>3234527	TACTCCTCCA	GGTAATGATG	AACCCTACAC	TGAGCAGATG	GGCAACTGTG
Consensus	TACTCCTCCA	GGTAATGATG	AACCCTACAC	TGAGCAGATG	GGCAACTGTG
>1737775IH	GAGAGAAGGG	TGAAAGGATC	CACCTCACTC	CTGATTTCAT	TGCAGGAAAA
>776455	GCGAG				
>3234527	GAGAGAAGGG	TGAAAGGATC	CACCTCACTC	CTGATTTCAT	TGCAGGAAAA
Consensus	GAGAGAAGGG	TGAAAGGATC	CACCTCACTC	CTGATTTCAT	TGCAGGAAAA
>1737775IH	AAGTTAGCTG	AATATGGACC	ACAAGGTAGG	GCATTTGTCC	ATGAGTGGGC
>3234527	AAGTTAGCTG	AATATGGACC	ACAAGGTAGG	GCATTTGTCC	ATGAGTGGGC
Consensus	AAGTTAGCTG	AATATGGACC	ACAAGGTAGG	GCATTTGTCC	ATGAGTGGGC
>1737775IH	TCATCTACGA	TGGGGAGTAT	TTGACGAGTA	CAATAATGAT	GAGAAATTCT
>3234527	TCATCTACGA	TGGGGAGTAT	TTGACGAGTA	CAATAATGAT	GAGAAATTCT
Consensus	TCATCTACGA	TGGGGAGTAT	TTGACGAGTA	CAATAATGAT	GAGAAATTCT

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>1737775IH >3234527	ACTTATCCAA ACTTATCCAA		CAAGCAGTAA	GATGTTCAGC	AGGTATTACT
Consensus			CAAGCAGTAA	GATGTTCAGC	AGGTATTACT
>1737775IH >1628724	GGTACAAATG	TAGTAAAGAA	GTGTCAGGGA	GGCAGCTGTT	ACACCAAAAG CAAAAG
Consensus	GGTACAAATG	TAGTAAAGAA	GTGTCAGGGA	GGCAGCTGTT	
>1737775IH				TGAAAAAGGA	
>1628724	ATGCACATTC	AATAAAGTAA	CAGGACTCTA	TGAAAAAGGA	TGTGAGTTTG
Consensus	ATGCACATTC	AATAAAGTAA	CAGGACTCTA	TGAAAAAGGA	TGTGAGTTTG
>1737775IH				CTATAATGTT	
>1628724	TTCTCCAATC	CCGCCAGACG	GAGAAGGCTT	CTATAATGTT	TGCACAACAT
Consensus	TTCTCCAATC	CCGCCAGACG	GAGAAGGCTT	CTATAATGTT	TGCACAACAT
>1737775IH	GTTGATTCTA	TAGTTGAATT	CTGTACAGAA	CAAAACCACA	ACAAAGAAGC
>1628724				CAAAACCACA	
>1427224	CTA	TAGTTGAATT	CTGTACAGAA	CAAAACCACA	ACAAAGAAGC
Consensus	GTTGATTCTA	TAGTTGAATT	CTGTACAGAA	CAAAACCACA	ACAAAGAAGC
>1737775IH	TCCAAACAAG	СААААТСААА	AATGCAATCT	CCGAAGCACA	TGGGAAGTGA
>1628724	TCCAAACAAG	CAAAATCAAA	AATGC		
>1427224	TCCAAACAAG	CAAAATCAAA	AATGCAATCT	CCGAAGCACA	TGGGAAGTGA
Consensus	TCCAAACAAG	СААААТСААА	AATGCAATCT	CCGAAGCACA	TGGGAAGTGA
>1737775IH	TCCGTGATTC	TGAGGACTTT	AAGAAAACCA	CTCCTATGAC	AACACAGCCA
>1427224	TCCGTGATTC	TGAGGACTTT	AAGAAAACCA	CTCCTATGAC	AACACAGCCA
Consensus	TCCGTGATTC	TGAGGACTTT	AAGAAAACCA	CTCCTATGAC	AACACAGCCA
>1737775IH	CCAAATCCCA	CCTTCTCATT	GCTGCAGATT	GGACAAAGAA	TTGTGTGTTT
>1427224	CCAAATCCCA	CCTTCTCATT	GCTGCAGATT	GGACAAAGAA	TTGTGTGTTT
>3235571				CAAAGAA	TTGTGTGTTT
Consensus	CCAAATCCCA	CCTTCTCATT	GCTGCAGATT	GGACAAAGAA	TTGTGTGTTT
>1737775IH	AGTCCTTGAC	AAATCTGGAA	GCATGGCGAC	TGGTAACCGC	CTCAATCGAC
>1427224	AGTCCTTGAC	AAATCTGGAA	GCATGGC		
>3235571	AGTCCTTGAC	AAATCTGGAA	GCATGGCGAC	TGGTAACCGC	CTCAATCGAC
Consensus				TGGTAACCGC	
>1737775IH	TGAATCAAGC	AGGCCAGCTT	TTCCTGCTGC	AGACAGTTGA	GCTGGGGTCC
>3235571	TGAATCAAGC	AGGCCAGCTT	TTCCTGCTGC	AGACAGTTGA	GCTGGGGTCC
Consensus					GCTGGGGTCC
>1737775IH	TGGGTTGGGA	TGGTGACATT	TGACAGTGCT	GCCCATGTAC	AAAGTGAACT
>3235571	TGGGTTGGGA	TGGTGACATT	TGACAGTGCT	GCCCATGTAC	AAAGTGAACT
Consensus	TGGGTTGGGA	TGGTGACATT	TGACAGTGCT	GCCCATGTAC	AAAGTGAACT

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>1737775IH >3235571	CATACAGATA	AACAGTGGCA	GTGACAGGGA	CACACTCGCC CACACTCGCC	AAAAGATTAC
Consensus	CATACAGATA	AACAGTGGCA	GTGACAGGGA	CACACTCGCC	AAAAGATTAC
>1737775IH				GCAGCGGGCT	TCGATCGGCA
>3235571			ACGTCCATCT		
Consensus	CTGCAGCAGC	TTCAGGAGGG	ACGTCCATCT	GCAGCGGGCT	TCGATCGGCA
>1737775IH	TTTACTGTGA	TTAGGAAGAA	ATATCCAACT	GATGGATCTG	AAATTGTGCT
>2680592		GAA	ATATCCAACT	GATGGATCTG	AAATTGTGCT
Consensus	TTTACTGTGA	TTAGGAAGAA	ATATCCAACT	GATGGATCTG	AAATTGTGCT
>1737775IH	GCTGACGGAT	GGGGAAGACA	ACACTATAAG	TGGGTGCTTT	AACGAGGTCA
>2680592	GCTGACGGAT	GGGGAAGACA	ACACTATAAG	TGGGTGCTTT	AACGAGGTCA
Consensus	GCTGACGGAT	GGGGAAGACA	ACACTATAAG	TGGGTGCTTT	AACGAGGTCA
>1737775ІН	AACAAAGTGG	TGCCATCATC	CACACAGTCG	CTTTGGGGCC	CTCTGCAGCT
>2680592	AACAAAGTNG	TGCCATCATC	CACACAGTCG	CTTTGGGGCC	CTCTGCAGCT
>608819	AACAAAGTGG	TGCCATCATC	CACACAGTCG	CTTTGGGGCC	CTCTGCAGCT
>608819IH	AACAAAGTGG	TGCCATCATC	CACACAGTCG	CTTTGGGGCC	CTCTGCAGCT
Consensus	AACAAAGTGG	TGCCATCATC	CACACAGTCG	CTTTGGGGCC	CTCTGCAGCT
>1737775IH	CAAGAACTAG	AGGAGCTGTC	CAAAATGACA	GGAGGTTTAC	AGACATATGC
>2680592	CAAGAACTAG	AGGAGCTGTC	CAAAATGACA	GGAGGTTTAC	AGACATATGC
>608819	CAAGAACTAG	AGGAGCTGTC	CAAAATGACA	GGAGGTTTAC	AGACATATGC
>608819IH	CAAGAACTAG	AGGAGCTGTC	CAAAATGACA	GGAGGTTTAC	AGACATATGC
Consensus	CAAGAACTAG	AGGAGCTGTC	CAAAATGACA	GGAGGTTTAC	AGACATATGC
>1737775IH	TTCAGATCAA	GTTCAGAACA	ATGGCCTCAT	TGATGCTTTT	GGGCCCTTT
>2680592	TTCAGATCAA	GTTCAGAACA	ATGGCCTCAT	TGATGCTTTT	GGGGCCCTTT
>608819	TTCAGATCAA	GTTCAGAACA	ATGGCCTCAT	TGATGCTTTT	GGGGCCCTTT
>608819IH	TTCAGATCAA	GTTCAGAACA	ATGGCCTCAT	TGATGCTTTT	GGGGCCCTTT
>1629002		GAACA	ATGGCCTCAT	TGATGCTTTT	GGGGCCCTTT
Consensus	TTCAGATCAA	GTTCAGAACA	ATGGCCTCAT	TGATGCTTTT	GGGGCCCTTT
>1737775IH	CATCAGGAAA	TGGAGCTGTC	TCTCAGCGCT	CCATCCAGCT	TGAGAGTAAG
>2680592	CATCAGGAAA	TGGAGCTGTC	TCTCAGCGCT	CCATCCAGCT	TGAGAGTAAG
>608819	CATCAGGAAA	TGGAGCTGTC	TCTCAGCGCT	CCATCCAGCT	TGAGAGTAAG
>608819IH	CATCAGGAAA	TGGAGCTGTC	TCTCAGCGCT	CCATCCAGCT	TGAGAGTAAG
>1629002		TGGAGCTGTC		CCATCCAGCT	
Consensus	CATCAGGAAA	TGGAGCTGTC	TCTCAGCGCT	CCATCCAGCT	TGAGAGTAAG

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Figure 1-D

>1737775IH >2680592	GGATTAACCC GGATTA	TCCAGAACAG	CCAGTGGATG	AATGGCACAG	TGATCGTGGA
>608819	GGATTAACCC	TCCAGAACAG	CCAGTGGATG	AATGGCACAG	TGATCGTGGA
>608819IH	GGATTAACCC	TCCAGAACAG	CCAGTGGATG	AATGGCACAG	TGATCGTGGA
>1629002	GGATTAACCC	TCCAGAACAG	CCAGTGGATG	AATGGCACAG	TGATCGTGGA
>1626435			•	GGCACAG	TGATCGTGGA
Consensus	GGATTAACCC	TCCAGAACAG	CCAGTGGATG	AATGGCACAG	TGATCGTGGA
>1737775ІН		GGAAAGGACA	CTTTGTTTCT	TATCACCTGG	ACAACGCAGC
>608819	CAG				
>608819IH			· -	TATCACCTGG	
>1629002				TATCACCTGG	
>1626435				TATCACCTGG	
Consensus	CAGCACCGTG	GGAAAGGACA	CTTTGTTTCT	TATCACCTGG	ACAACGCAGC
>1737775IH	CTCCCCAAAT	CCTTCTCTGG	GATCCCAGTG	GACAGAAGCA	AGGTGGCTTT
>608819IH	CTCCCCAAAT	CCTTCTCTGG	GATCCCAGTG	GACAGAAGCA	AGGTGGCTTT
>1629002	CTCCCCAAAT	CCTTCTCTGG	GATCCCAGTG	GAC	
>1626435	CTCCCCAAAT	CCTTCTCTGG	GATCCCAGTG	GACAGAAGCA	AGGTGGCTTT
>1582693					GGCTTT
Consensus	CTCCCCAAAT	CCTTCTCTGG	GATCCCAGTG	GACAGAAGCA	AGGTGGCTTT
>1737775IH				CTCCAAATCC	
>608819IH	GTAGTGGACA	AAAACACCAA	AATGGCCTAC	CTCCAAATCC	CAGGCATTGC
>1626435				CTCCAAATCC	
>1582693	GTAGTGGACA	AAAACACCAA	AATGGCCTAC	CTCCAAATCC	CAGGCATTGC
Consensus	GTAGTGGACA	AAAACACCAA	AATGGCCTAC	CTCCAAATCC	CAGGCATTGC
>1737775IH				AGCAAGCTCA	
>608819IH				AGCAAGCTCA	
>1626435	*			AGCAAGCTCA	
>1582693				AGCAAGCTCA	
Consensus	TAAGGTTGGC	ACTTGGAAAT	ACAGTCTGCA	AGCAAGCTCA	CAAACCTTGA
>1737775IH ·	CCCTGACTGT	CACGTCCCGT	GCGTCCAATG	CTACCCTGCC	TCCAATTACA
>608819IH	CCCTGACTGT	CACGTCCCGT	GCGTCCAATG	CTACCCTGCC	TCCAATTACA
>1626435	CCCTGACTGT	CA			
>1582693	CCCTGACTGT	CACGTCCCGT	GCGTCCAATG	CTACCCTGCC	TCCAATTACA
Consensus	CCCTGACTGT	CACGTCCCGT	GCGTCCAATG	CTACCCTGCC	TCCAATTACA
>1737775IH				AAATTCCCCA	
>608819IH				AAATTCCCCA	GCCCTCTGGT
>1582693	GTGACTTCCA	AAACGAACAA	GGACACCAGC		
>1281379				AAATTCCCCA	
Consensus	GTGACTTCCA	AAACGAACAA	GGACACCAGC	AAATTCCCCA	GCCCTCTGGT

DOLLESS. OSETOS



Figure 1-E

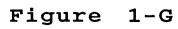
>1737775IH	AGTTTATGCA	AATATTCGCC	AAGGAGCCTC	CCCAATTCTC	AGGGCCAGTG
>608819IH	AGTTTATGCA	AATATTCGCC	AAGGAGCCTC	CCCAATTCTC	AGGGCCAGTG
>1281379	AGTTTATGCA	AATATTCGCC	AAGGAGCCTC	CCCAATTCTC	AGGGCCAGTG
Consensus	AGTTTATGCA	AATATTCGCC	AAGGAGCCTC	CCCAATTCTC	AGGGCCAGTG
>1737775IH	TCACAGCCCT	GATTGAATCA	GTGAATGGAA	AAACAGTTAC	CTTGGAACTA
>608819IH	TCACAGCCCT	GATTGAATCA	GTGAATGGAA	AAACAGTTAC	CTTGGAACTA
>1281379	TCACAGCCCT	GATTGAATCA	GTGAATGGAA	AAACAGTTAC	CTTGGAACTA
Consensus	TCACAGCCCT	GATTGAATCA	GTGAATGGAA	AAACAGTTAC	CTTGGAACTA
>1737775IH			TGATGCTACT		
>608819IH			TGATGCTACT		
>1281379	CTGGATAATG	GAGCAGGTGC	TGATGCTACT	AAGGATGACG	GTGTCTACTC
Consensus	CTGGATAATG	GAGCAGGTGC	TGATGCTACT	AAGGATGACG	GTGTCTACTC
>1737775IH			ACACGAATGG		
>608819IH			ACACGAATGG		
>1281379	AAGGTATTTC	ACAACTTATG	ACACGAATGG	TAGATACAGT	
>1628509					GTGC
Consensus	AAGGTATTTC	ACAACTTATG	ACACGAATGG	TAGATACAGT	GTAAAAGTGC
>1737775IH	CCCCTCTCCC	አርር አርሞሞል እር	GCAGCCAGAC	CGAGAGTGAT	ACCCCAGCAG
>608819IH			GCAGCCAGAC		
>1281379	GGGCTC	HOOHOTTANC	ocnocenone	001.01.010111	
>1628509		አርር አርጥጥል እር	GCAGCCAGAC	ССУСУСТСУТ	ACCCCAGCAG
Consensus			GCAGCCAGAC		
Consensus	GGGCTCTGGG	AGGAGTTAAC	GCAGCCAGAC	GGAGAGIGAI	
>1737775IH	AGTGGAGCAC	TGTACATACC	TGGCTGGATT	GAGAATGATG	AAATACAATG
>608819IH	AGTGGAGCAC	TGTACATACC	TGGCTGGATT	GAGAATGATG	AAATACAATG
>1628509	AGTGGAGCAC	TGTACATACC	TGGCTGGATT	GAGAATGATG	AAATACAATG
Consensus	AGTGGAGCAC	TGTACATACC	TGGCTGGATT	GAGAATGATG	AAATACAATG
		•			
>1737775IH			TTAATAAGGA		
>608819IH			TTAATAAGGA		
>1628509	GAATCCACCA	AGACCTGAAA	TTAATAAGGA	TGATGTTCAA	CACAAGCAAG
Consensus	GAATCCACCA	AGACCTGAAA	TTAATAAGGA	TGATGTTCAA	CACAAGCAAG
>1737775IH			TCGGGAGGCT		
>608819IH					TTCTGATGTC
>1628509			TCGGGAGGCT		
Consensus	TGTGTTTCAG	CAGAACATCC	TCGGGAGGCT	CATTTGTGGC	TTCTGATGTC
>1737775IH	CC 3 3 3 TC CTC	CCMMACCMCA	ጥርጥርጥጥርርር	ССТССССААА	TCACCGACCT
>1/3///SIH >608819IH					TCACCGACCT
>6088191H >1628509		CCATACCTGA		CCIGGCCAAA	ICACCGACCI
	CCAAATGCTC	CCATACCIGA	101		ACCT
>2060739	CC	ርር እጠእርር መ ር እ	መርመርመመርርር እ	CCTCCCAAA	TCACCGACCT
Consensus	CCAAATGCTC	CCATACCIGA	TOTOTTCCCA	CCIGGCCMAA	ICACCGACCI

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Figure 1-F

>1737775IH	CAACCCCCAA	ATTCACCCC	CCACTCTCAT	TAATCTGACT	TCCACACCTC
>608819IH				TAATCTGACT	
>2060739				TAATCTGACT	
Consensus				TAATCTGACT	
			001.01010111		100110110010
>1737775IH	CTGGGGATGA	TTATGACCAT	GGAACAGCTC	ACAAGTATAT	CATTCGAATA
>608819IH	CTGGGGATGA	TTATGACCAT	GGAACAGCTC	ACAAGTATAT	CATTCGAATA
>2060739	CTGGGGATGA	TTATGACCAT	GGAACAGCTC	ACAAGTATAT	CATTCGAATA
Consensus	CTGGGGATGA	TTATGACCAT	GGAACAGCTC	ACAAGTATAT	CATTCGAATA
>1737775IH	AGTACAAGTA	TTCTTGATCT	CAGAGACAAG	TTCAATGAAT	CTCTTCAAGT
>608819IH	AGTACAAGTA	TTCTTGATCT	CAGAGACAAG	TTCAATGAAT	CTCTTCAAGT
>2060739	AGTACAAGTA	TTCTTGATCT	CAGAGACAAG	TTCAATGAAT	CTCTTCAAGT
Consensus	AGTACAAGTA	TTCTTGATCT	CAGAGACAAG	TTCAATGAAT	CTCTTCAAGT
>1737775IH				CAACTCTGAG	
>608819IH				CAACTCTGAG	
>2060739				CAACTCTGAG	
Consensus	GAATACTACT	GCTCTCATCC	CAAAGGAAGC	CAACTCTGAG	GAAGTCTTTT
4505555					
>1737775IH				ATGGCACAGA	
>608819IH				ATGGCACAGA	
>2060739	TGTTTAAACC	AGAAAACA'I'I'	AC'I"I"I"I'GAAA	ATGGCACAGA	
>1624934					TCTTTTCATT
Consensus	TGTTTAAACC	AGAAAACA'I'T	AC'I"I"I"I'GAAA	ATGGCACAGA	TCT"I"I"TCATT
>1737775IH	CCTATTCACC	COCOMO A TA A	CCTCCATCTC	AAATCAGAAA	татссаасат
>608819IH				AAATCAGAAA	
>1624934				AAATCAGAAA	
Consensus				AAATCAGAAA	
Consensus	GCIATICAGG	CIGIIGAIAA	GGICGAICIG	AMICAGAM	TATCCAMCAT
>1737775IH	TGCACGAGTA	TCTTTGTTTA	TTCCTCCACA	GACTCCGCCA	GAGACACCTA
>608819IH	TGCACGAGTA	TCTTTGTTTA	TTCCTCCACA	GACTCCGCCA	GAGACACCTA
>1624934	TGCACGAGTA	TCTTTGTTTA	TTCCTCCACA	GACTCCGCCA	GAGACACCTA
>1800229		GTTTA	TTCCTCCACA	GACTCCGCCA	GAGACACCTA
Consensus	TGCACGAGTA	TCTTTGTTTA	TTCCTCCACA	GACTCCGCCA	GAGACACCTA
>1737775IH	GTCCTGATGA	AACGTCTGCT	CCTTGTCCTA	ATATTCATAT	CAACAGCACC
>608819IH	GTCCTGATGA	AACGTCTGCT	CCTTGTCCTA	ATATTCATAT	CAACAGCACC
>1624934		AACGTCTGCT			
>1800229	GTCCTGATGA	AACGTCTGCT	CCTTGTCCTA	ATATTCATAT	CAACAGCACC
Consensus	GTCCTGATGA	AACGTCTGCT	CCTTGTCCTA	ATATTCATAT	CAACAGCACC
>1737775IH				TGGAAGTGGA	
>608819IH				TGGAAGTGGA	
>1800229				TGGAAGTGGA	
>990829					TAGGAGAACT
Consensus	ATTCCTGGCA	TTCACATTTT	AAAAATTATG	TGGAAGTGGA	TAGGAGAACT

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>1737775IH >608819IH >1800229 >990829 Consensus	GCAGCTGTCA GCAGCTGTCA GCAGCTGTCA	ATAGCCTAGG ATAGCCTAGG ATAGCCTAGG ATAGCCTAGG	GCTGAATTTT GCTGAATTTT GCTGAATTTT GCTGAATTTT GCTGAATTTT	TGTCAGATAA TGTCAGATAA	АТАААТАА АТААААТАА АТААААТАА АТААААТАА АТААААТАА
>1737775IH	TCATTCATCC				
>608819IH	TCATTCATCC	TTTTTTTGA	TTATAAAATT	TTCTAAAATG	TATTTTAGAC
>1800229	TCATTCATCC	TTTTTTTGA	TTATAAAATT	TTCTAAAATG	TATTTTAGAC
>990829	TCATTCATCC	TT			
Consensus	TCATTCATCC	TTTTTTTGA	TTATAAAATT	TTCTAAAATG	TATITTAGAC
>608819IH >1800229	TTCCTGTAGG TTCCTGT	GGGCGATATA	CTAAATGTAT	ATAGTACATT	TATACTAAAT
Consensus	TTCCTGTAGG	GGGCGATATA	CTAAATGTAT	ATAGTACATT	ТАТАСТАААТ
>608819IH	GTATTCCTGT	AGGGGGCGAT	ATACTAAATG	TATTTTAGAC	TTCCTGTAGG
Consensus	GTATTCCTGT	AGGGGGCGAT	ATACTAAATG	TATTTTAGAC	TTCCTGTAGG
>608819IH	GGGCGATAAA				
Consensus	GGGCGATAAA	ATAAAATGCT	AAACAACTGG	GTA	

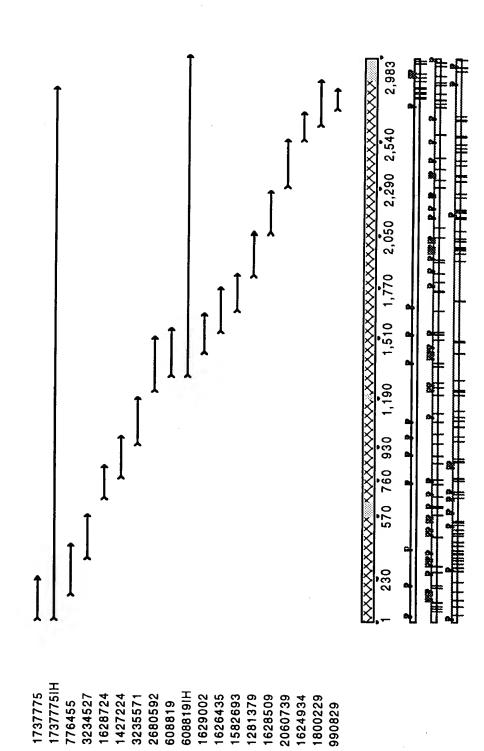
~ Figure

> 1628724 3234527

776455

1427224 3235571

DSO4555.03E798



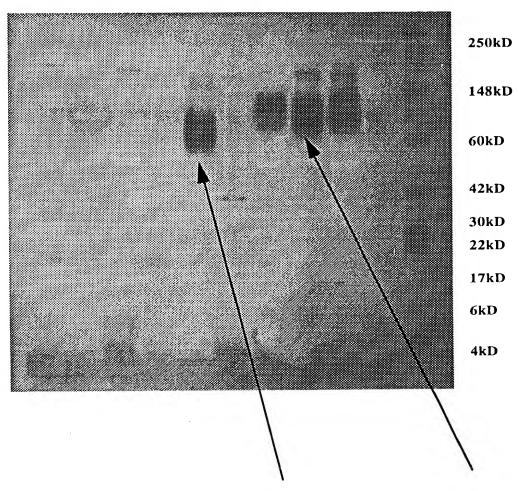
1582693 1281379 1628509 2060739 1624934 1800229

990829

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Figure 3

1 2 3 4 5 6 7 8 9 10 11



Lane	Tissue	Lane	Tissue
1	Prostate	7	Colon
2	Breast	8	Colon
3	Bladder	9	Colon
4	Lung	10	Blank
5	Colon cancer	11	Markers
6	Colon cancer		

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